

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/646,390A

Source: FFWO

Date Processed by STIC: 11-15-04

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IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/646,390A

DATE: 11/15/2004

TIME: 12:00:48

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\11152004\J646390A.raw

(pg. 6)

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3 <110> APPLICANT: SAITOU, Mitinori
4   SURANI, Azim
6 <120> TITLE OF INVENTION: Genes
8 <130> FILE REFERENCE: 674558-2002.1
10 <140> CURRENT APPLICATION NUMBER: 10/646,390A
11 <141> CURRENT FILING DATE: 2003-08-21
13 <150> PRIOR APPLICATION NUMBER: 10/621,911
14 <151> PRIOR FILING DATE: 2003-07-17
16 <150> PRIOR APPLICATION NUMBER: PCT/GB02/00215
17 <151> PRIOR FILING DATE: 2002-01-18
19 <150> PRIOR APPLICATION NUMBER: GB 0101300.2
20 <151> PRIOR FILING DATE: 2001-01-18
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27 <211> LENGTH: 617
28 <212> TYPE: DNA
29 <213> ORGANISM: Mus musculus
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34 caaactacga aagaatcaag gaagaatatg aggtggctga gatgggggca ccgcacggat 180
35 cggcttctgt cagaactact gtgatcaaca tgcccagaga ggtgtcgggt cctgaccatg 240
36 tggctctggtc cctgttcaat acactcttca tgaacttctg ctgcctgggc ttcatagcct 300
37 atgcctactc cctgaagtct agggatcgga agatgggtgg tgatgtgact ggagcccagg 360
38 cctacgcctc cactgctaag tgcctgaaca tcagcacctt ggctcctcagc atcctgatgg 420
39 ttgttatcac cattgttagt gtcacatca ttgtttcttaa cgctcaaaac cttcacactt 480
40 aatagaggat tccgacttcc ggtcctgaag tgcttcaccc tccgcagctg cgtccctcct 540
41 tgcccctccc tacacgcagg tgtaacactc atttatctat ccacagtgga ttcaataaag 600
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45 <211> LENGTH: 137
46 <212> TYPE: PRT
47 <213> ORGANISM: Mus musculus
49 <400> SEQUENCE: 2
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51 1          5          10          15
53 Pro Pro Asn Tyr Glu Arg Ile Lys Glu Tyr Glu Val Ala Glu Met
54          20          25          30
56 Gly Ala Pro His Gly Ser Ala Ser Val Arg Thr Thr Val Ile Asn Met
57          35          40          45
59 Pro Arg Glu Val Ser Val Pro Asp His Val Val Trp Ser Leu Phe Asn
60          50          55          60

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62 Thr Leu Phe Met Asn Phe Cys Cys Leu Gly Phe Ile Ala Tyr Ala Tyr
63 65              70              75              80
65 Ser Val Lys Ser Arg Asp Arg Lys Met Val Gly Asp Val Thr Gly Ala
66              85              90              95
68 Gln Ala Tyr Ala Ser Thr Ala Lys Cys Leu Asn Ile Ser Thr Leu Val
69              100             105             110
71 Leu Ser Ile Leu Met Val Val Ile Thr Ile Val Ser Val Ile Ile Ile
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85 agaaagtcga cccaatgaag gacctgaaa ctcctcagaa gaaagatgaa gaggacgctt 180
86 tggatgatac agacgtccta caaccagaaa cactagtaaa ggtcatgaaa aagctaacct 240
87 taaaccccg tgtcaagcgg tccgcacgcc ggcgcagtct acggaaccgc attgcagccg 300
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92 aaatcgggaa gaattaggag cttacattgt acgctgccct ggctgtcgac gatgccgcac 600
93 agcagatgtg aaagctatct tttgtttaag attaaacttt ttctggtgct gggaaatctt 660
94 aacttggtta cctttaaatt gtagatagga tgcacaacga tccagattta tgtgaagttt 720
95 agaagcctca agctgtgagg cccagggctg aggaataaag taaatagaat ttggagtatg 780
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111              35              40              45
113 Gly Val Lys Arg Ser Ala Arg Arg Arg Ser Leu Arg Asn Arg Ile Ala
114              50              55              60
116 Ala Val Pro Val Glu Asn Lys Ser Glu Lys Ile Arg Arg Glu Val Gln
117 65              70              75              80
119 Ser Ala Phe Pro Lys Arg Arg Val Arg Thr Leu Leu Ser Val Leu Lys
120              85              90              95
122 Asp Pro Ile Ala Lys Met Arg Arg Leu Val Arg Ile Glu Gln Arg Gln
123              100             105             110
125 Lys Arg Leu Glu Gly Asn Glu Phe Glu Arg Asp Ser Glu Pro Phe Arg

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136 <212> TYPE: DNA
137 <213> ORGANISM: Rattus sp
139 <400> SEQUENCE: 5
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142 cgggggtttt ggaagcccg cggccctctc ttctcacctt gctccactag cacgcggtg 180
143 ttttctactga gccagcact ggctaagtgg agcaccagga gtttcaggct atccttcaga 240
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150 cccaaatcct aggattaaag gtttttagtg gctgcagtgt tgggtggggg ttgttttgat 660
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182 catttatttt tggattttca gactccctcc ccatagtcca aactggccct ccagttcagt 2580
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190 acatttcttt tgagttgcaa tatcctgatt aacatttttc tttcagagac gagatgagcc 3060
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225 <211> LENGTH: 1444
226 <212> TYPE: DNA
227 <213> ORGANISM: Rattus sp
229 <400> SEQUENCE: 6

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232 aaaaaacgag tccccatctc caatggctcg ttctaactcg ttctgtgtat ttattaaaga 180
233 taacaaattht gcctctatta caaatthtct tgcaaaactag aaaatctaaa ataaaagatc 240
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235 aatctgtcat tccccatgaat ccaaatgtca cacatcctat ataattttaa ggtagcaag 360
236 tagagatttc cccagaccca agaaaagttt aatcttaaac aaaaaaacag ctttcacatc 420
237 tgctgcatgg caccgttaac ggcaaggaca gtgtatgatt caaactgcc taattcttct 480
238 ggttctgccc aatttttagca ttctcagaag gatcccatct ctgataatgg cagaaagtac 540
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246 caaataaaat aaaatcaccg cccaacccca cccctataaa aaaccgcgcg accacaccac 1020
247 atatactccc cccccccgc accatcacta catcacctc tccacctatt cccacctccc 1080
248 cccccaacat taaccccacc ccatcaggga aaccccacac accaacaat aaattagaca 1140
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252 ccaattaaac acgcccacc caccacaaca cccaccgccc tgacaacaca cttcggaact 1380
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259 <213> ORGANISM: Rattus sp

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263 <222> LOCATION: (7471)..(7471)

264 <223> OTHER INFORMATION: "n" is an unknown nucleotide

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267 <221> NAME/KEY: misc_feature

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269 <223> OTHER INFORMATION: "n" is an unknown nucleotide

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272 <221> NAME/KEY: misc_feature

273 <222> LOCATION: (7608)..(7608)

274 <223> OTHER INFORMATION: "n" is an unknown nucleotide

276 <400> SEQUENCE: 7

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279 atgctggaga aagaattcaa aagttaggta gagaactctg ttgagcagag cacttgcttt 180
280 tcttcagag gatctgagtt caagtccag gacctatct acagttttct gtaactctag 240
281 ctccagaggg tctgacactt ctgttcactg tgggcacctg cattcacaga caaacataaa 300
282 gtagtctatc acccttttca cagaaaaccc acagcatgtg aggaaatccg ggtctctgcg 360
283 caatgcccc acagcagaag gggggagctg gagagatggt tcatctgtta gcccatatat 420

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:7; N Pos. 7471,7554,7608

Seq#:8; N Pos. 2115,2142,2143,2146

VERIFICATION SUMMARY

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L:401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:7440
M:341 Repeated in SeqNo=7
L:467 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:2100